



Deep-sea benthic habitats modeling and mapping in a NE Atlantic seamount (Galicia Bank)



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ABSTRACT

This study presents the results of seafloor habitat identification and mapping of a NE Atlantic deep seamount. An “assemble first, predict later” approach has been followed to identify and map the benthic habitats of the Galicia Bank (NW Iberian). Biotic patterns inferred from the survey data have been used to drive the definition of benthic assemblages using multivariate tools. Eight assemblages, four hard substrates and four sedimentary ones, have been described from a matrix of structural species. Distribution of these assemblages was correlated with environmental factors (multibeam and backscatter data) using binomial GAMs. Finally, the distribution model of each assemblage was applied to produce continuous maps and pooled in a final map with the distribution of the main benthic habitats. Depth and substrate type are key factors when determining soft bottom communities, whereas rocky habitat distribution is mainly explained by rock slope and orientation. Enrichment by northern water masses (LSW) arriving to GB and possible zooplankton biomass increase at vertical-steep walls by “bottom trapping” can explain the higher diversity of habitat providing filter-feeders at slope rocky breaks. These results concerning vulnerable species and habitats, such as *Lophelia* and *Madrepora* communities and black and bamboo coral aggregations were the basis of the Spanish proposal of inclusion within the Natura 2000 network. The aim of the present study was to establish the scientific criteria needed for managing and protecting those environmental values.

1. Introduction

There is a global call to protect marine species and ecosystems and to develop coherent and connected networks of Marine Protected Areas (MPAs) in national and international waters (Bullimore et al., 2013; Edgar et al., 2014). The last Convention on Biological Diversity (CBD) established in 1992, states the necessity of conserving 10% of the coastal and marine areas through effective and equitably managed, ecologically representative, and well-connected systems of protected areas and other effective area-based conservation measures. Similar calls have been made by regional organisations, such as the Oslo-Paris (OSPAR) Convention and several countries around the world which are contributing to this global effort by developing their own MPA networks (IUCN and UNEP-WCMC, 2013). In Europe, the Marine Strategy Framework Directive (MSFD) includes a requirement to establish an ecologically coherent network of MPAs to help protect vulnerable

species and habitats. Despite the remarkable expansion of the number of MPAs and the amount of marine areas protected in recent years, MPAs still cover only 2.8% of the world's seas and oceans (IUCN and UNEP-WCMC, 2013), far from the 10% target for 2020. As part of the necessary effort to conserve marine ecosystems, the Spanish government has proposed eleven off-shore areas as Special Areas of Conservation (SACs) under the Habitat Directive (Council Directive 92/43/EEC). These areas were shortlisted based on previous knowledge and their priority features studied within the research project INDEMARES (EC contract LIFE 07/ NAT /E/000732). The main objective of this project was to provide the necessary information to establish a network of representative MPAs in Spanish waters.

Habitat identification and mapping are fundamental in determining the locations for potential protected areas although both requirements are challenging, especially in the deep sea (Bryan and Metaxas, 2007; Howell, 2010; Howell et al., 2010; González-Mirelis et al., 2012;

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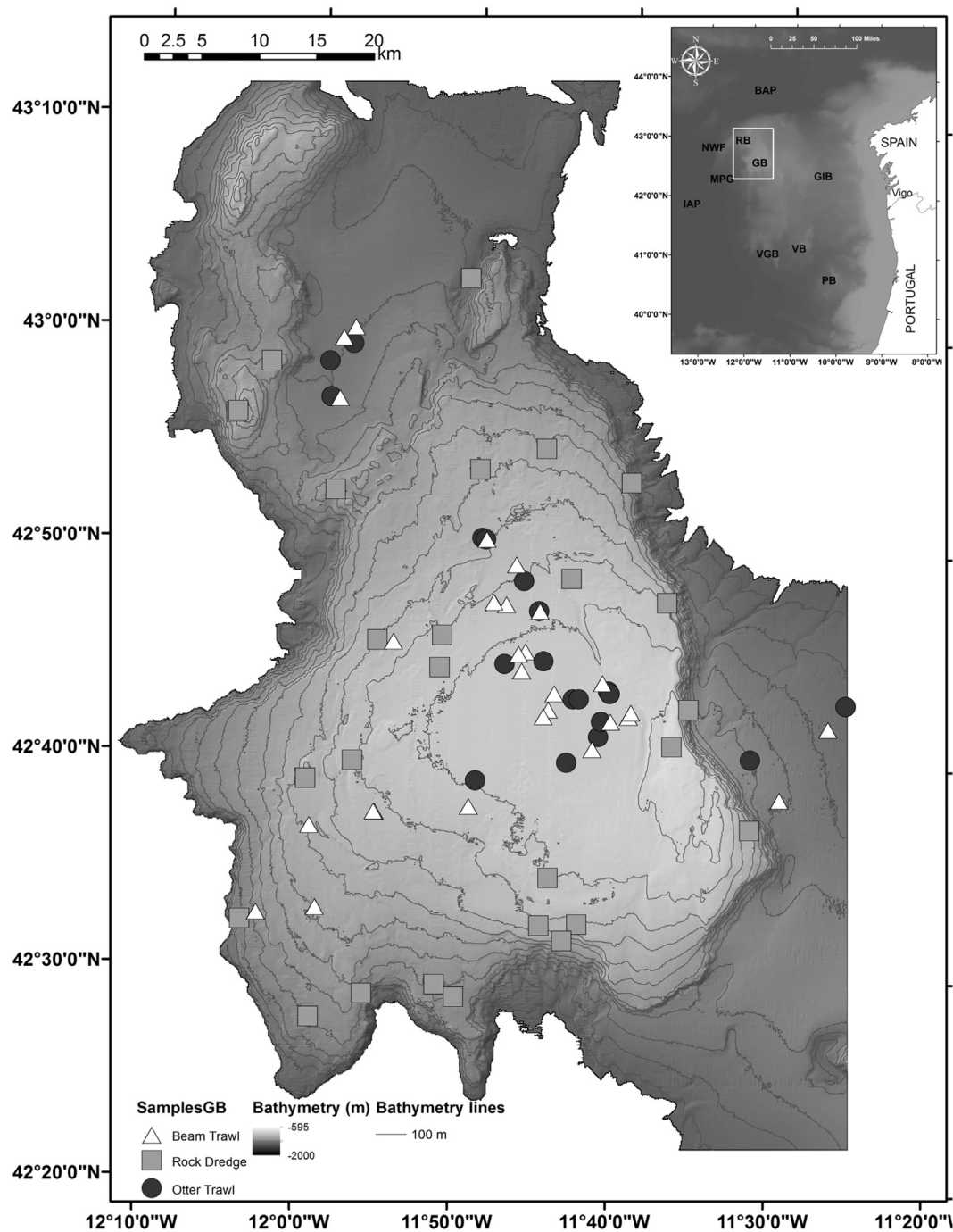


Fig. 1. Study area and sampling stations. Bathymetric contours each 100 m. Inset: Location of the study area on the west Iberia continental margin. GB, Galicia Bank; RB, Rucabado Bank, VGB, Vasco da Gama bank; VB, Vigo bank; PB, Porto bank; GIB, Galicia Interior Basin; TZ, Transition Zone; NWF, Northwestern flank; DGM, Deep Galicia Margin; BAP, Biscay abyssal plain; IAP, Iberia abyssal plain. Bathymetry from the Spanish EEZ Project (multibeam bathymetry) and the GEBCO Digital Atlas.

Bullimore et al., 2013; Davies et al., 2015; González-Irusta et al., 2015). Deep-sea ecosystems are usually characterised by their remoteness and challenging accessibility. Great depths mean expensive costs in time and budget. Moreover, whereas in the shallow-water ecosystems the benthic assemblages are relatively well known, the description of these communities in the deep sea is still poor (Bullimore et al., 2013). Furthermore, hard-bottom living communities (typified mainly by corals and sponges), due to their patchy, irregular distribution linked to remote bottom accidents such as canyons, seamounts, steep reliefs, etc., are even more inaccessible than soft-bottom communities in the deep sea, and consequently information on their distribution and zonation is comparatively scarce. Therefore, it is necessary to describe these

assemblages, especially hard-bottom ones, and the underlying environmental conditions which structure them in order to identify the habitats to protect (Howell, 2010; Howell et al., 2010). Only after properly describing the habitats it is possible to include them in the different habitat classification systems, a required step to protect them through directives (Habitats Directive- HD, Marine Strategy Framework Directive- MSFD) and conventions (OSPAR). EUNIS habitat classification (Davies and Moss, 2002) was used to classify benthic habitats and to extend crosswalks to the above cited directives and conventions and in order to supply information that can be used in the SAC management plan.

Once the different habitats have been identified, it is necessary to

map their extent. The distribution of these habitats based on empirical studies offers a discrete and incomplete view of their distribution (Greathead et al., 2014). Distribution models (DM), also called habitat suitability models, species DM or ecological niche models, are correlative approaches that use discrete distribution data and full spatial coverage of environmental data to explain and predict patterns of distribution (Elith and Graham, 2009; Reiss et al., 2015). Although these models have been used mainly to predict species distribution (e.g. Bryan and Metaxas, 2007; Monk et al., 2010; González-Irusta et al., 2015), they have also been used to predict the distribution of habitats: essential fish habitats (González-Irusta and Wright, 2016), habitat mapping based on habitat-forming species distribution (Howell et al., 2011), and benthic communities defined by visual criteria (Martín-García et al., 2013) or multivariate analysis (Moritz et al., 2013).

The Galicia Bank (GB) is the deepest SAC of the Spanish proposal. This area was considered relevant for further research in the framework of the INDEMARES project because of the presence of well developed and conserved priority habitats such as *Lophelia* and *Madrepora* communities and black and bamboo coral aggregations (Duineveld et al., 2004; Somoza et al., 2014), of which some are catalogued as vulnerable (OSPAR, 2008; Annex I Habitat Directive, 92/43/EEC). Vulnerable species such as deep-water sharks (Piñeiro et al., 2001; Bañón et al., 2006, 2008; Rodríguez-Cabello et al., 2014) and carnivorous sponges (Cristobo et al., 2015) inhabit GB as well. Moreover, GB is one of the few seamounts in the Spanish Economic Exclusive Zone (EEZ). There are at least some 800 major seamounts in the North Atlantic, nevertheless most of them occur in high seas, associated with the Arctic Mid-Ocean Ridge, the Mid-Atlantic Ridge, and the Greenland-Iceland/Iceland-Faeroe Rise, all of which are large features that dominate the topography of the seabed (Gubbay, 2003). GB is not that type of seamount, showing several singularities derived from its proximity to the continent, its deep summit (Surugiu et al., 2008) and its location on the confluence of water masses and currents (Cartes et al., 2014).

In this study an “assemble first, predict later” approach (González-Mirelis et al., 2012; Robert et al., 2016) has been followed to identify and map the benthic habitats of the GB. Biotic patterns inferred from the survey data have been used to drive the definition of benthic assemblages using multivariate tools. This multivariate analysis was used to explore the composition and distribution of different epibenthic assemblages observed on the GB. After classifying the assemblage of each sample, their distribution was correlated with environmental factors using binomial General Additive Models (GAMs) in a DM framework. Finally, the DMs of each assemblage were applied to produce continuous maps with the location of the main habitats and pooled in a final map with the distribution of the main benthic habitats of GB.

2. Material and methods

2.1. Study area

GB is an isolated non-volcanic seamount (Black et al., 1964), located 120 nautical miles west of the NW Spanish shoreline (Fig. 1). The outline of its summit is almost triangular-shaped, being 75 km long in a NNE-SSW direction, by 58 km wide in a WNW-ESE direction, and with a total surface of 1844 km². Across its summit surface, depths range from 600 m (to the SE) to more than 2000 m (to the W). To the east, GB borders on a transitional zone of the seamounts' domain leading to the Galicia Interior Basin; to the N-NW, Rucabado Bank is located, connecting with the Galicia escarpment (Vanney et al., 1979), named the Northwestern Flank by Vázquez et al. (2008); to the W-SW, Deep Galicia Margin (Murillas et al., 1990) is found; S-SE, the seamounts domain extends, connecting GB with other seamounts, such as the Vasco da Gama, Vigo and Porto banks, as well as other minor hills (Fig. 1).

Surrounding GB, at depths sampled in this study (to 1800 m), we intercepted 3 water masses direct/indirectly related with the communities sampled: i) the Eastern North Atlantic Central Water (ENACW)

occupies the level of hydrographic structures found over the GB summit; ii) Below ENACW and to ca. 1500 m, we found the Mediterranean Outflow Water (MOW) characterised by an increase of salinity. MOW had a maximum core, with higher salinity and low-oxygen concentration at depths between 800 and 1100–1200 m (Ambar and Howe, 1979; Iorga and Lozier, 1999; INDEMARES data). This water mass comes from the Strait of Gibraltar, and reaches velocities of 5–10 cm/s (Iorga and Lozier, 1999), and iii) below MOW, the Labrador Sea Water (LSW) that moves southwards from northern latitudes to GB and has a core at 1800 m (Iorga and Lozier, 1999).

2.2. Biological samples

Habitats and species data were obtained from three multi-disciplinary surveys performed on the GB in the summer of 2009, 2010 and 2011 (Ecomarg0709; BanGal0810; BanGal0811). Mega-epibenthic fauna were collected using two different sampling systems, beam trawl and otter trawl (GOC73), in sedimentary grounds, and a rock dredge in rocky grounds. The beam trawl used consists of two steel skids joined by a 3.5 m horizontal bar with a vertical opening of 1 m and mesh size of 10 mm. Trawling was carried out during 15 min at an average speed of 2 knots. The otter trawl used was a 2-warp trawl with two doors, 35.7 m headline length, 30 m bridles, a vertical height or opening of 2.7 m and 20 mm mesh at the codend. Haul duration was 45 min at 2.8–2.9 knots. Trawl openings were recorded by means of SCANMAR and Simrad ITI Trawl monitoring sensors mounted in the mouths of the trawls. Rock dredge consists in a heavy frame of 80 × 30 cm with a 10 mm mesh size net. Haul duration was 5 min at 1.5 knots.

Otter trawl and beam trawl faunal data is quantitative and expressed in biomass (wet weight) whereas rock dredge faunal data was standardised as biomass percentage of each sample. During the three surveys, a total of 27 rock dredges, 19 otter trawls and 29 beam trawls were carried out across the seamount from 737 to 1809 m deep (Fig. 1).

2.3. Environmental layers

The multibeam bathymetric and backscatter data used during the present investigation were collected in the frame of the Spanish EEZ Project, during the ZEE-2001, ZEE-2002, ZEE-2003 and ZEE-2007 cruises carried out aboard the R.V. Hespérides. The multibeam echosounder systems used were a Kongsberg-Simrad EM-12 (ZEE-2001 to ZEE-2003 cruises), operating 81 beams at a 12 kHz frequency, and a Kongsberg-Simrad EM-120 (ZEE-2007 cruise), operating 191 beams at a 12 kHz frequency. Simultaneously to the multibeam data, information on the shallow structure of the area was obtained with a hull-mounted TOPAS PS 018 high-resolution parametric profiler system. Multibeam data set was processed using Caris Hips & Sips V.7.1 software and interpolated to a 75 × 75 m sided regular grid, providing virtually continuous coverage of the entire survey area. The processed bathymetry (supplementary Fig. 1a) was used to produce four additional variables of the seafloor using the spatial analyst and the Benthic Terrain Model tools (ARCGIS 9.3), namely; northness (supplementary Fig. 1b) and eastness (supplementary Fig. 1c), slope (supplementary Fig. 1d) and fine Bathymetric Position Index (BPI, supplementary Fig. 1e). The multibeam backscatter was also processed with the Geocoder module of the Caris Hips & Sips software.

The backscatter intensity is the energy reflected and represents the roughness and hardness of the substratum (supplementary Fig. 2). Sediments were collected with a USNEL box corer (Hessler and Jumars, 1974). Particle size analysis was performed by a combination of dry sieving and sedimentation techniques (Buchanan, 1984). Geomorphologic interpretation, seafloor type interpretation and granulometry were combined to produce a new layer with three different substrate types (supplementary Fig. 1f).

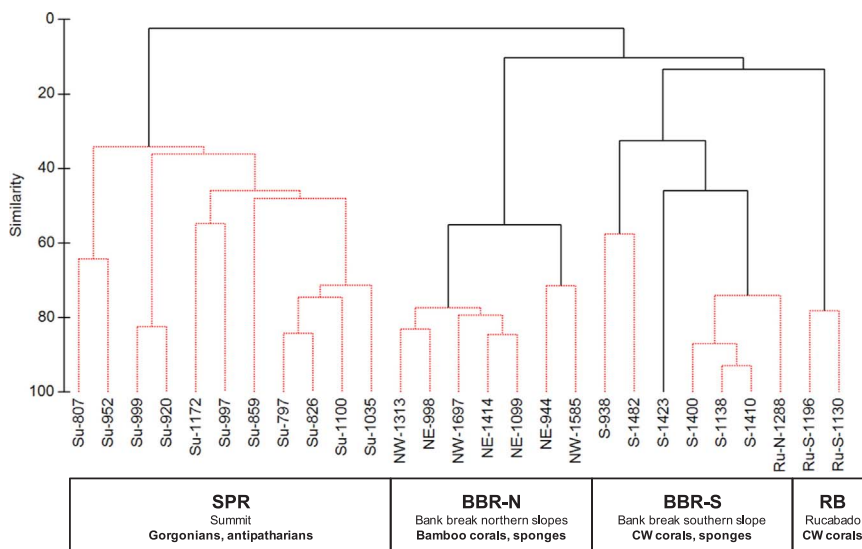


Fig. 2. Cluster analysis for structural species in rock dredges. Solid black lines indicate significant evidence of structure (SIMPROF test, $P > 0.05$). Dotted red lines indicate no evidence of structure. Labels sign GB zone/slope (Su: summit plain rock; NW-NE-S: Northwestern, Northeastern, Southern slopes; Ru: Rucabado Bank, N-north and S-south) and depth (m). Assemblages are defined by GB zone/slope, and SIMPER more typificant species (CW: cold-water).

2.4. Data analysis

Habitat identification and mapping of the biological communities followed a two step procedure (Moritz et al., 2013). First, the structural species assemblages were identified using clustering analysis. In a second step, distribution of the assemblages in the GB was predicted using binomial GAM models in a DM framework. Cluster and SIMPER analyses were conducted using PRIMER 6, whereas modeling was performed using R 3.02 (R Core Team, 2013).

In order to describe the structural species assemblages, biological data from otter trawl and beam trawl data were pooled together (after standardization to a common area). Trawl and dredge matrices were reduced, considering only structural species, defined as sessile, three-dimensional, large-bodied (mainly cnidarians and sponges), or those accompanying megafauna which appear in large numbers, with a limited motility, such as echinoderms or crustaceans. Benthic assemblages have been analysed determining groups of samples by means of hierarchical cluster analysis performed on log-transformed otter and beam trawl biomass matrix, and on rock dredge biomass percentages data matrix, using the Bray–Curtis similarity index. A similarity profile (SIMPROF) test was used as a prerequisite for defining assemblages (Clarke et al., 2008). SIMPER analyses were also run to identify the species typifying assemblages.

Once the structural species assemblages were identified, they were mapped using DMs. The only exception was the RB assemblage. This assemblage only had two presence points and therefore it was not suitable for modeling. The RB presence points were joined with BBR-S for mapping purposes. The DMs use different mathematical algorithms to calculate the ecological niche of the target species based on the environmental variable values at the presence point (Monk et al., 2010). The assemblage's probability of presence was modelled using binomial GAMs, with logit as link function, which is a technique that performed well when compared with others (González-Irujo et al., 2015). The implementation gam in the package “mgcv” (Wood, 2011) was used. To avoid overfitting all the smoothers were constrained to 4 knots, limiting degrees of freedom to a maximum of 3. For the variable selection, an Akaike information criterion (AIC)-based stepwise procedure was used in both directions for the GAM (Akaike, 1973). The full binomial model for all the habitats was:

$$Pp = \beta_1 + s(\text{bathymetry}) + s(\text{northness}) + s(\text{depth}) + s(\text{eastness}) + s(\text{slope}) + s(\text{BPI}) + f(\text{bottom type}) + \epsilon_1 \quad (1)$$

Where Pp is the probability of presence for each habitat, β is the

intercept, s is an isotropic smoothing function (thin plate regression splines, one for each variable and model), f indicates the variable which was included as factor in the formula and ϵ is the error term. Before starting the analysis, the correlation between the explanatory variables was checked for collinearity using Spearman rank correlations and Variance Inflation Factors (VIFs) (Zuur et al., 2009). Spearman Rank values were lower than 0.5 and the VIFs lower than 3 so all the variables were included in the model. The spatial autocorrelation of residuals was tested visually using variograms, with implementation vario in the package “gstat” (Pebesma, 2004).

The performance of the models was tested using cross-validation. The presence-absence data for each assemblage was randomly divided into a training subsample (with 80% of the total points) and a test subsample (with the other 20%). The ability of the training subsample to predict the probability of presence was tested using the test subsample. The performance of the models was estimated using two different statistics: the Area under the Curve (AUC) of the receiver operating characteristic (ROC, Fielding and Bell, 1997) and the kappa statistic (Cohen, 1960). The process was repeated 10 times for each combination of species and model, calculating the AUC and Kappa values each time based on a different random selection of training and test subsample. Both statistics were calculated using the implementation of *evaluate* in the R package ‘dismo’. The threshold used to compute the kappa value was calculated each time, using *threshold* in the same package. The threshold that provided maximum kappa values was applied to produce the final communities map. Finally, in order to show a unique map with all the communities together, we merged them selecting for each pixel the community with the highest probability of presence. Previously, all the values lower than its habitat's prevalence were replaced by 0. Pixels which 0 values for all the habitats selected were classified as “no prediction”.

3. Results

3.1. Assemblages identification

Similarity between rocky samples is shown in Fig. 2. Clustering pattern is not determined by depth like the first factor, but for a combination of slope and orientation. Cluster groups are typified by the combination of structural taxa as are cold-water corals, bamboo corals, black corals, gorgonians and large sponges.

Four main clusters can be defined at a distance cut of 20 (significant at SIMPROF test $P > 0.05$), corresponding with four assemblages of characteristic typificant fauna and environmental variables:

Table 1
SIMPER results for rocky habitats. Species typifying cluster groups.

SPR: Summit plain rock		BBR-N: Bank break N-E-W slopes	
Average similarity: 44.0	Sim %	Average similarity: 67.3	Sim %
<i>Acanthogorgia armata</i>	35.6	<i>Acanella arbuscula</i>	72.5
<i>Parantipathes</i> sp.	21.1	Hexactinellida	10.8
<i>Narella bellissima</i>	10.2	<i>Aphrocallistes beatrix</i>	5.9
<i>Swiftia rosea</i>	5.8	Geodiidae	4.8
<i>Trissopathes</i> sp.	4.3	<i>Anthothela grandiflora</i>	3.2
<i>Schizopathes</i> sp.	1.9	<i>Phakellia robusta</i>	1.9
<i>Bathypathes</i> sp.	1.5	<i>Brisinga endecacnemus</i>	0.9
BBR-S: Bank break S slopes		RB: Rucabado Bank	
Average similarity: 50.2	Sim %	Average similarity: 78.2	Sim %
<i>Madrepora oculata</i>	35.4	<i>Madrepora oculata</i>	48.5
<i>Lophelia pertusa</i>	24.2	<i>Lophelia pertusa</i>	40.2
Hexactinellida	8.1	<i>Desmophyllum dianthus</i>	3.5
<i>Aphrocallistes beatrix</i>	5.4	<i>Trissopathes</i> sp.	0.3
<i>Asconema setubalense</i>	4.1	<i>Leiopathes</i> sp.	0.2
<i>Phakellia robusta</i>	1.0	<i>Asconema setubalense</i>	0.2
<i>Desmophyllum dianthus</i>	0.6	<i>Aphrocallistes beatrix</i>	0.1

- i) **SPR (Summit Plain Rock)**: typified by the gorgonian *Acanthogorgia armata* and the black coral *Parantipathes* sp., together with other gorgonians such as *Swiftia rosea* and *Narella bellissima*, and antipatharians such as *Trissopathes* sp. and *Schizopathes* sp. (Table 1).
- ii) **BBR-N (Bank Break Rock)**: located mainly in the northern, western and eastern slopes of the bank. Typified by the bamboo coral *Acanella arbuscula*, and with a lower contribution several sponge species (Hexactinellida undet., *Aphrocallistes beatrix*, Geodiidae undet., *Phakellia robusta*), gorgonians (*Anthothela grandiflora*), and the sea star *Brisinga endecacnemus*.
- iii) **BBR-S (southern Bank Break Rock)**: located in the southern slope of the bank. With the presence of colonies of cold-water corals (*Madrepora oculata*, *Lophelia pertusa*), and a companion fauna of species than also typified BBR-N (large sponges, the bamboo coral *Acanella arbuscula*) and cold-water coral epifauna as solitary scleractinian corals (*Desmophyllum dianthus*).
- iv) **Rucabado Bank (RB)**: Cold-water corals reef (*Madrepora oculata*, *Lophelia pertusa*) and their epibiont fauna (*Desmophyllum dianthus*, black corals) and large sponges (*Asconema setubalense*, *Aphrocallistes beatrix*).

Although depth was not the main factor structuring hard-bottom communities, some depth-related trends were identified. SPR (split in the first dichotomy) occupied shallower depths (797–1172 m, mean = 951 ± 47 m) than BBR-N (944–1697 m, mean = 1293 ± 291 m), BBR-S (938–1482 m, mean = 1297 ± 195 m) and RB (1130–1196 m, mean = 1163 ± 47 m), seeing significant differences among assemblages (Kruskal-Wallis test: Chi-sq. = 11.4; $p = 0.01$) and (Bonferroni test, paired comparisons) between SPR-BBR-S ($p = 0.009$) and SPR-RB depths ($p = 0.008$).

Differences in habitat complexity are shown in Fig. 3. This figure was computed using the average richness of structural (tridimensional) species: 15.1 ± 3.2 (mean \pm SD) in BBR-S, 14.5 ± 3.6 in RB, 8.9 ± 2.5 in BBR-N, and 6.0 ± 2.7 in SPR. Therefore, there are significant differences between BBR-S and RB assemblages and BBR-N and SPR assemblages. The most complex assemblages, BBR-S and RB, are characterised by colonial scleractinians (*Lophelia* and *Madrepora*, Table 2) whereas the simplest one, SPR, are characterised by gorgonians and antipatharians. Gorgonians and black corals did not present clear differences among assemblages, whereas hexactinellids, demosponges and bamboo corals present a higher occurrence in the bank break than in the summit plain rock. Finally, colonial scleractinians are more frequent in the southern slope and in the Rucabado Bank (Fig. 3).

In the soft bottoms, the dendrogram analysis (Fig. 4) clearly shows the existence of three main clusters (at 18% similarity). One of these

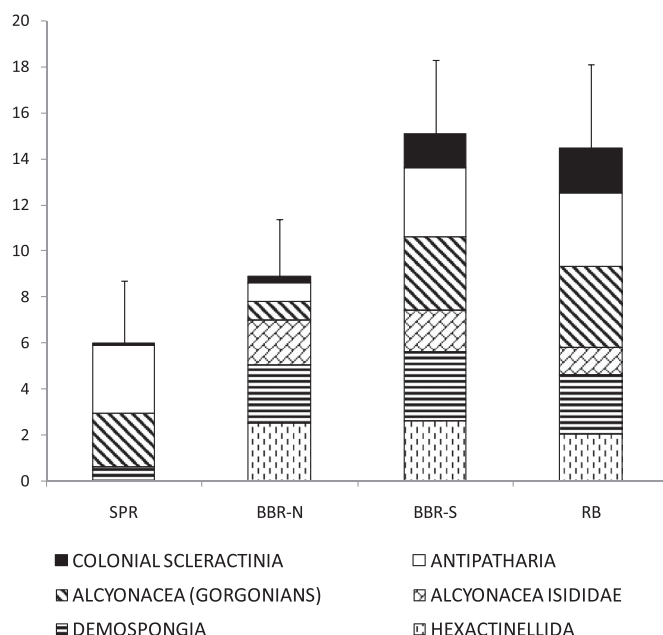


Fig. 3. Mean three-dimensional sessile large species richness by rocky assemblage. Bars represent standard error.

Table 2
Species typifying 75% of intragroup similarity.

SS- Bank summit (750–780 m)		SSrf (780–1000 m)	
Average similarity: 37.4	Sim %	Average similarity: 51.3	Sim %
Ophiacanthidae	33.3	<i>Lophelia pertusa</i>	16.0
<i>Flabellum chunii</i>	12.2	<i>Madrepora oculata</i>	11.8
<i>Deltocyathus eccentricus</i>	10.9	<i>Urothychus</i> spp.	10.1
<i>Cancer bellianus</i>	5.8	<i>Acanthogorgia</i> spp.	8.9
<i>Limopsis</i> spp.	5.7	<i>Desmophyllum cristagali</i>	8.1
<i>Thenea muricata</i>	4.2	<i>Munidopsis</i> spp.	7.6
<i>Chaceon affinis</i>	4.1	Geodiidae indet.	5.4
		<i>Parantipathes</i> spp.	4.7
		<i>Lima marioni</i>	3.3
BBS- Bank break (1000–1200 m)		FS- Bank flanks (1400–1800 m)	
Average similarity: 30.35	Sim %	Average similarity: 33.3	Cum %
<i>Cidaris cidaris</i>	23.2	<i>Benthogone rosea</i>	21.3
<i>Thenea muricata</i>	21.5	<i>Neolithodes grimaldii</i>	10.9
<i>Peltaster placenta</i>	17.6	<i>Araeosoma fenestratum</i>	10.5
<i>Acanella arbuscula</i>	9.5	<i>Glyphocrangon longirostris</i>	5.9
<i>Colus</i> spp.	2.3	<i>Colossendeis colosseae</i>	5.8
<i>Chaceon affinis</i>	1.7	<i>Stephanocyathus</i> sp.	4.4
		<i>Fissidentallium capillosum</i>	4.2
		<i>Umbellula</i> sp.	3.8
		<i>Peltaster placenta</i>	3.7
		<i>Acanella arbuscula</i>	3.6
		<i>Swiftia rosea</i>	3.1

groups includes all the summit samples and was also divided into two subgroups. These clusters and subclusters have been confirmed by SIMPROF ($P < 0.05$) and were considered faunal assemblages. Main factors explaining assemblages are depth and depth-related factors as substrate type and water masses. Gear used to obtain samples was not a decisive factor.

SIMPER analysis (Table 2) highlight intergroup faunal differences. The 4 assemblages identified for sedimentary habitats are:

- i) **Summit Sands (SS)**: shallowest samples (750–780 m), substrate of medium sands of low reflectivity, affected by the ENACW. Sands dwelled by a huge abundance (9.5 kg/ha) of ophiuroids of the

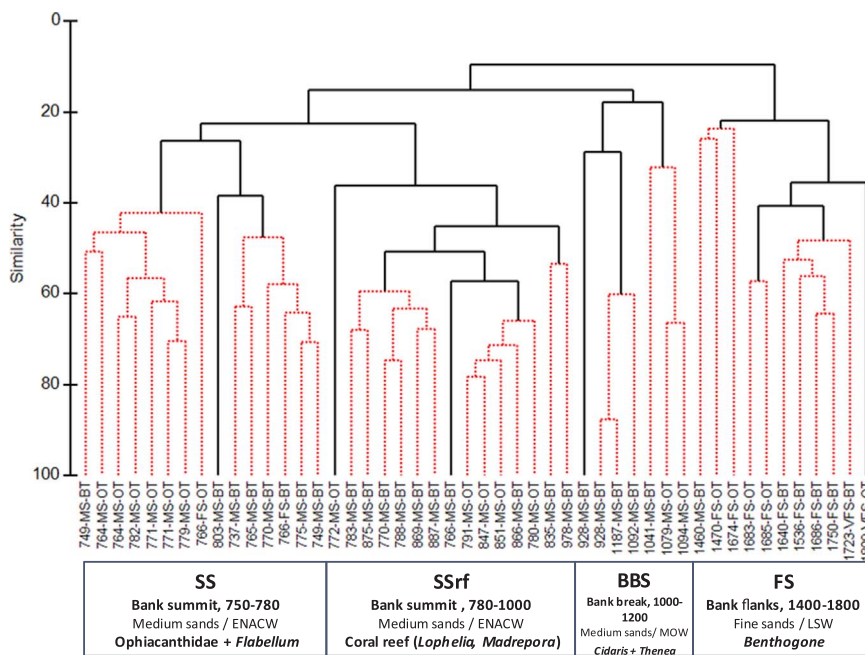


Fig. 4. Cluster analysis for structural species in sedimentary areas. Solid lines indicate significant evidence of structure (SIMPPOF test, $P > 0.05$). Dotted lines indicate no evidence of structure. Labels sign depth (m), sediment type (MS- Medium sands, FS- Fine sands, VFS: very fine sands), and gear (BT- beam trawl, OT- otter trawl). Assemblages are defined by depth, substrate type, water mass and SIMPER more typificant species.

family Ophiacanthidae (*Ophiacantha* sp.) and Ophiolidae (*Ophiomyces grandis*), and with lower densities of the solitary coral *Deltocyathus eccentricus* and *Flabellum chunii*, and the bivalve *Limopsis* spp. (*L. minuta* and *L. cristata*). Companion mobile fauna is characterised by the crab *Cancer bellianus* and *Chaceon affinis*. This cluster has an internal separation of otter trawl and beam trawl samples, at 25% similarity (Fig. 1), due to the different catchability of both samplers to small species including ophiuroids and *D. eccentricus*.

- ii) **Summit Sands with CW coral reef patches (SSrf):** Substrate of medium sands of low backscatter with depths ranging from 780 to 1000 m, also in the ENACW layer. Sands covered by patches of cold-water coral colonies (*Lophelia pertusa* and *Madrepora oculata*, 3.3 and 3.2 kg/ha of live coral). Together with these two scleractinian species, typifying species are mostly species living on coral such as *Desmophyllum cristagalli*, *Acanthogorgia* spp., *Lima marioni*, *Parantipathes* sp., and epibiont fauna such as different species of *Munidopsis* and *Urophychus*.
- iii) **Bank Break Sands (BBS):** substrate of medium sands of medium reflectivity located in depths ranging from 1000 to 1200 m, in the MOW layer. Sands dominated by the pencil urchin *Cidarid cidaris*, the sponge *Thenea muricata*, the sea star *Peltaster placenta*, the gastropod *Colus* spp. and the bamboo coral *Acanella arbuscula* (also abundant in rocky habitats).
- iv) **Bank Flanks Sands (FS):** fine and very fine sands of low reflectivity in depths ranging from 1400 to 1800 m, LSW mass. Clearly dominated by the holothurian *Elasipodida Benthogone rosea* (6.3 kg/ha), accompanied by the leather urchin *Araeosoma fenestratum*, a very distinct arthropod fauna (the crab *Neolithodes grimaldii*, *Glyphocrangon longirostris* or the giant sea spider *Colossendeis colossae*), and several anthozoa including *Umbellula* sp., *Acanella arbuscula* and *Swiftia rosea*.

3.2. Habitat classification

These habitats characterised by clustering have to be included in EUNIS (European Union Nature Information System) habitat classification system (Davies and Moss, 2002) and related with the catalogues and annexes of European directives and conventions.

Table 3 summarises correspondences and links among habitat described in the present paper, EUNIS levels and OSPAR List of

Threatened and/or Declining Species and Habitats (OSPAR, 2008) and Habitat Directive Annex I (92/43/EEC). Cold-water coral habitats classification scheme recently developed by Davies et al. (2017) was also included. Most of GB habitats formed by corals can be included in one or more than one categories of this scheme (Table 3). The only exception is the absence of hard substrate habitat types formed by the bamboo coral *Acanella arbuscula*, common in the GB rocky grounds.

Habitats described in this paper were assigned to HD Annex 1 (Table 3) when they fulfill a density criterium. Hence, only BBR, BBRS, and SSrf were assigned to the HD habitat type 1170 due to their density, and also their structural complexity.

3.3. Habitat mapping

The distribution of the biological communities was mapped using binomial GAMs. Model performance was good in all cases with high values of explained deviance, AUC and kappa values (Table 4).

Sediment type and depth were included in all the models highlighting the importance of these variables defining biological assemblages. Slope was only included in the rock assemblage SPR whereas the orientation of the slope (eastness or northness) was included in the other two rock communities (BBR-N and BBR-S-RB) from soft and hard bottoms; SSrf, BBR and BBRS. The different soft communities showed clear bathymetry segregation, with a low level of overlapping except at the summit of the seamount (Fig. 5). In this area, SSrf and SS overlap in depths of around 800 m although with a clearly different distribution, deeper for SSrf and shallower for SS. The BBS assemblage occupied an intermediate depth between the summit and the deepest areas of the seamount which were covered by the FS assemblage. In the case of the rock communities there was also a bathymetric distribution although slope was the main environmental driver in the distribution of the communities. The flat rock bottoms were occupied by the SPR assemblage whereas the rock areas with higher slopes were occupied by BBR-N (mainly in slopes with north-west orientation) and BBRS (mainly in the seamount slopes with a southern orientation). SPR showed a preference for shallower rocky areas than the other two rock assemblages. BBR-S had a peak in the probability of presence around 1200 m whereas BBR-N had the highest values in the probability of presence in the deeper areas.

Model maps per habitat were merged in a unique map selecting for each pixel the habitat with the highest probability of presence (Fig. 6)

Table 3
Habitat characterisation at different levels and links with OSPAR list of threatened habitats and Annex I of Habitat Directive.

GB goohabitat	EUNIS 3	GB habitats	EUNIS 4–6	Davies et al. (2017)	OSPAR list	HD
Plain rock (summit)	A6.1 Deep-sea rock A6.2 Deep-sea mixed substrata A6.6 Deep-sea bioherms	Summit plain rock with gorgonians and black corals	A6.11 Deep-sea bedrock A6.13 Deep-sea manganese nodules A6.722 Summit communities of seamount within the mesopelagic zone	6 CW Antipatharians and/or Gorgonians on Hard Substrate 6.1.1 Antipatharians on Hard Substrate 7.1.1 Isolated colonies of Scleractinians, Antipatharians and Gorgonians on Hard/Mixed Substrate or Consolidated Mud	Coral garden Deep-sea aggregations	1170 sponge
Steep rock (bank break and slope)	A6.1 Deep-sea rock A6.7 Raised features of the deep-sea bed A6.1 Deep-sea rock A6.7 Raised features of the deep-sea bed	Bank break rock with bamboo corals, gorgonians and large sponges Bank break rock with cold-water coral colonies, black & bamboo corals, and large sponges	A6.11 Deep-sea bedrock A6.14 Boulders on the deep-sea bed A6.62 Deep-sea sponge aggregations A6.621 Facies with <i>Phoronema grayi</i> A6.11 Deep-sea bedrock A6.61 Communities of deep-sea corals A6.62 Deep-sea sponge aggregations A6.14 Boulders on the deep-sea bed A6.22 Deep-sea biogenic debris A6.75 Carbonate mounds	6.2 CW Gorgonians on Hard/Mixed Substrate or Compact Mud 3.3.3 Isolated Colonies of <i>Madrepora oculata</i> and <i>Lophelia pertusa</i> on Hard Substrate 8.1.4 Antipatharians, Short Sponges and Sparse Large Sponges on Hard Substrate	Coral garden Deep-sea aggregations Lophelia reefs Coral garden Deep-sea aggregations Carbonate mounds	1170 sponge
Medium sands (summit)	A6.1 Deep-sea rock A6.6 Deep-sea bioherms A6.7 Raised features of the deep-sea bed A6.3 Deep-sea sand A6.2 Deep-sea mixed substrata A6.3 Deep-sea sand A6.6 Deep-sea bioherms	Cold-water corals reef Summit medium sands with Ophiacanthidae and <i>Flabellum chunii</i> Summit medium sands with cold-water corals reef patches	A6.11 Deep-sea bedrock A6.61 Communities of deep-sea corals A6.611 Deep-sea <i>Lophelia pertusa</i> reefs A6.722 Summit communities of seamount within the mesopelagic zone A6.61 Communities of deep-sea corals A6.611 Deep-sea <i>Lophelia pertusa</i> reefs A6.722 Summit communities of seamount within the mesopelagic zone A6.75 Carbonate mounds	1.1.3 Mixed <i>Madrepora oculata</i> and <i>Lophelia pertusa</i> reef 1.2.2 CW Scleractinian Reef Colonised by Antipatharians and/or Gorgonians 10.1.2 Flabellidae on Soft Substrate	Lophelia reefs Carbonate mounds Coral garden	1170
Fine and very fine sands (flanks)	A6.2 Deep-sea mixed substrata A6.3 Deep-sea sand A6.4 Deep-sea muddy sand	Bank break medium sands with <i>Cidaris</i> and <i>Thenea muricata</i> Bank flanks fine sands with elasipodid holothurians (<i>B. rosea</i>)	A6.22 Deep-sea biogenic debris A6.75 Carbonate mounds A6.722 Summit communities of seamount within the mesopelagic zone A6.724 Flanks of seamount or bank	9.1.1 Isolated Colonies of <i>Lophelia pertusa</i> and <i>Madrepora oculata</i> on Soft Substrate 1.4.2 Loosely-packed <i>Lophelia pertusa</i> and/or <i>Madrepora oculata</i> Framework with Soft Substrate Colonised by Antipatharians	Lophelia reefs Carbonate mounds	1170

Table 4
Summary of GAM results. Assemblage label from Figs. 2 and 3. AUC: Area Under the Curve.

Assemblage	GAM Formula	Explained deviance	AUC	Kappa
SS	$P_p = \beta_1 + s(\text{depth}) + f(\text{sediment}) + \varepsilon_1$	65.2%	0.95 ± 0.02	0.84 ± 0.08
SSrf	$P_p = \beta_2 + s(\text{depth}) + s(\text{eastness}) + f(\text{sediment}) + \varepsilon_2$	54.4%	0.86 ± 0.04	0.64 ± 0.05
BS	$P_p = \beta_3 + s(\text{depth}) + f(\text{sediment}) + \varepsilon_3$	99.8%	0.99 ± 0.01	0.94 ± 0.08
FS	$P_p = \beta_4 + s(\text{depth}) + f(\text{sediment}) + \varepsilon_4$	100%	1	1
SPR	$P_p = \beta_5 + s(\text{depth}) + s(\text{slope}) + f(\text{sediment}) + \varepsilon_5$	71.4%	0.94 ± 0.11	0.8 ± 0.31
BBR-N	$P_p = \beta_6 + s(\text{depth}) + s(\text{eastness}) + s(\text{northness}) + f(\text{sediment}) + \varepsilon_6$	60%	0.86 ± 0.13	0.6 ± 0.35
BBR-S-RB	$P_p = \beta_7 + s(\text{depth}) + s(\text{northness}) + f(\text{sediment}) + \varepsilon_7$	77.6%	0.80 ± 0.12	0.55 ± 0.12

after replacing by 0 all the values lower than prevalence. The pattern of habitat distribution in this probability map match with the patterns described in the cluster analysis (Fig. 2, Fig. 4), with depth (water masses) and substrate type as key factors in sedimentary areas and slope orientation in rocky areas. Some habitats have a clear geographical location in cluster analysis (Fig. 2), but modelling predicts their distribution in a wider area. This is the case of BBR-S, cluster of the southern slope and Rucabado Bank samples, with a higher probability of presence all along the bank (Fig. 6).

4. Discussion

Epibenthic assemblages from Galicia Bank have a discernible ecological structure which can be considered a proxy for benthic habitats. These habitats have been inferred from the assemblages obtained in

dendrograms through the environmental interpretation and modeling. Depth and substrate type (and also the depth-related influence of water masses) were key factors in sedimentary habitats whereas rocky habitats were determined by slope and slope orientation, although with certain level of bathymetry segregation as well. Eight habitats have been described, 4 on hard substrates and 4 on sedimentary ones.

4.1. Bathymetrical zonation of GB habitats

In most seamount studies, depth was the strongest environmental proxy for the assemblage-structuring processes (McClain and Lundsten, 2014; Du Preez et al., 2016). This depth-regulated environment generates communities which were generally distributed as bands encircling the seamount (Du Preez et al., 2016). These bands are present in the GB, even though depth-related zonation was clearer on

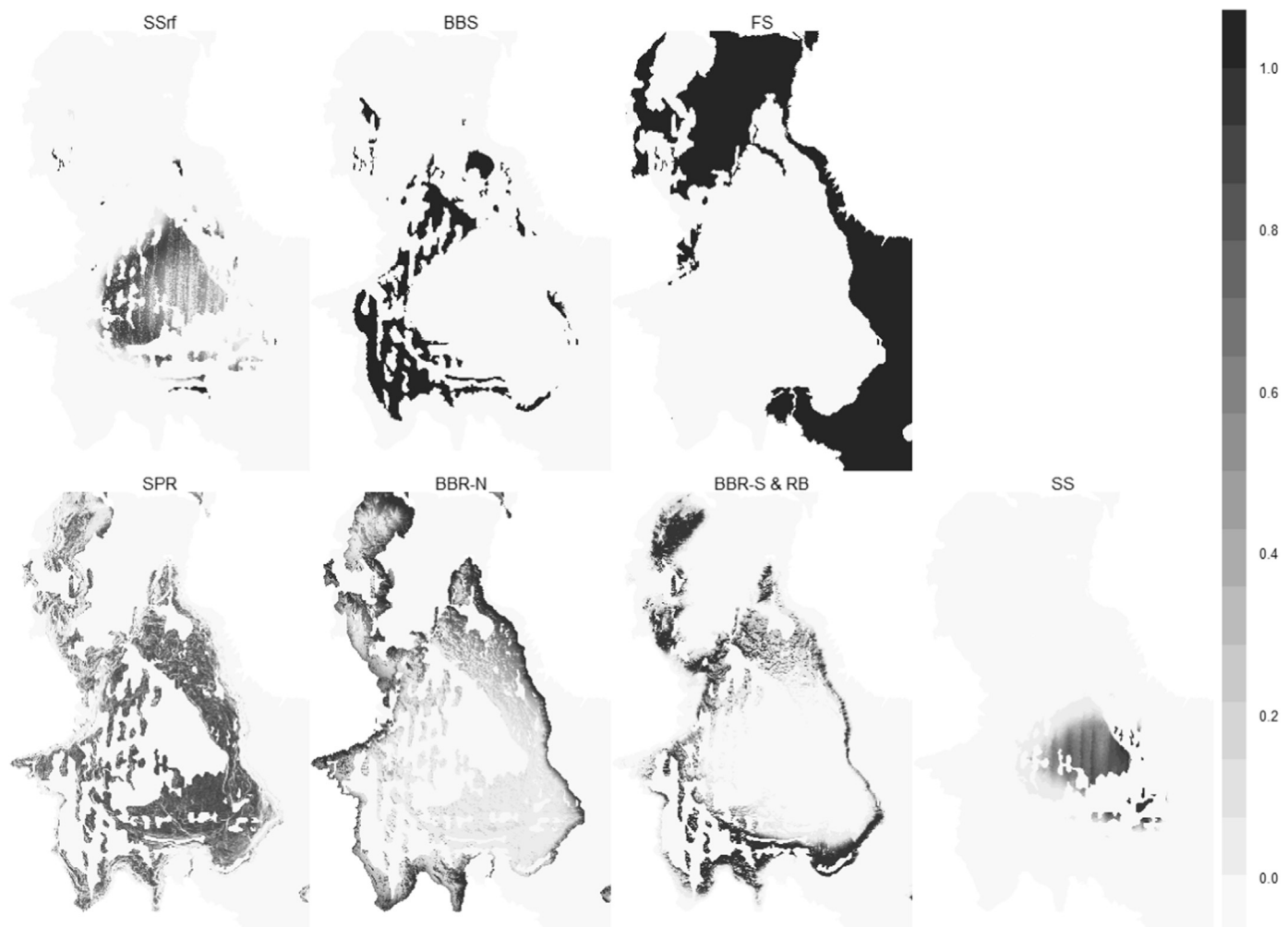


Fig. 5. Maps of probability of presence of every GB habitats.

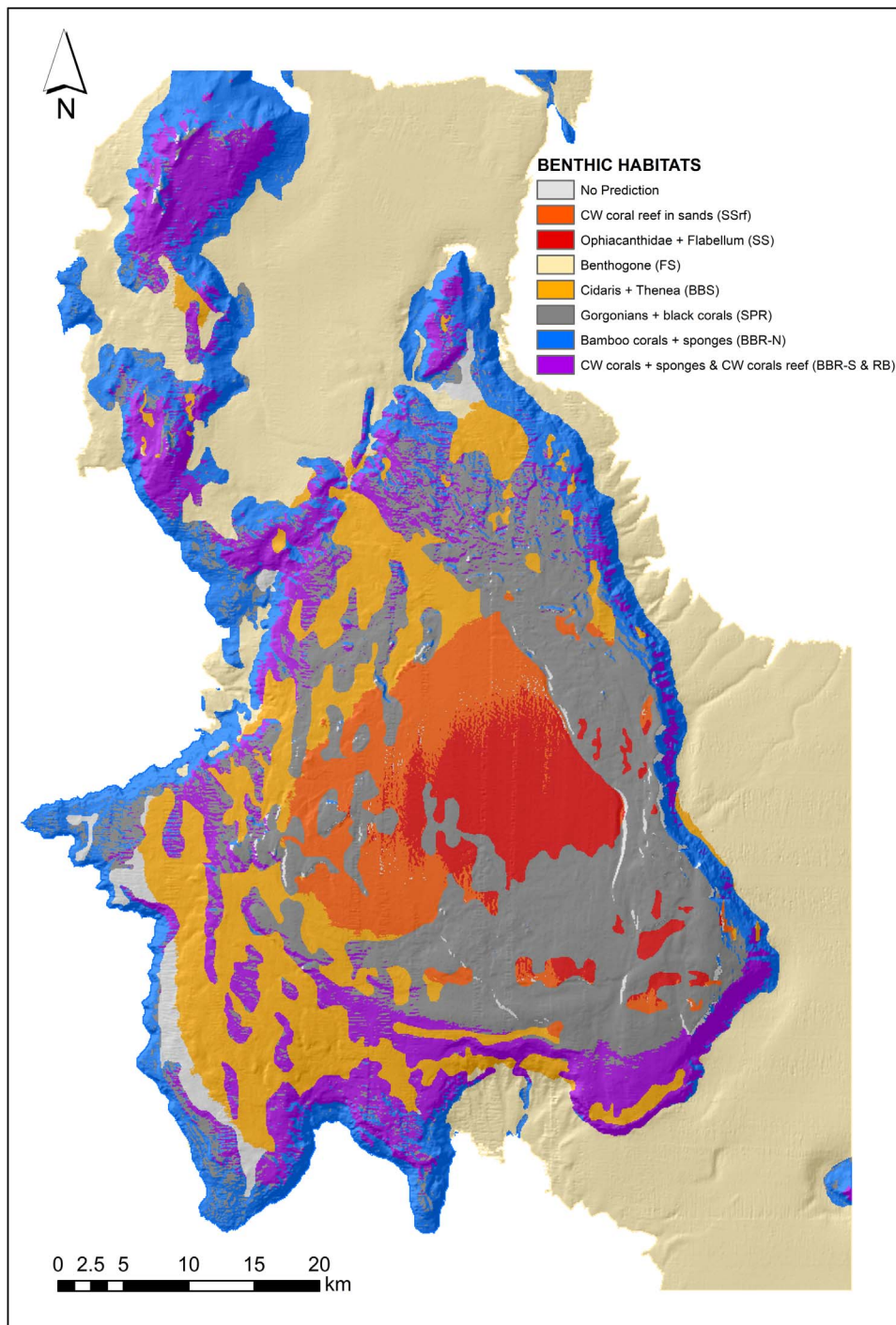


Fig. 6. Habitat map (highest probability of presence).

sedimentary than in rocky benthic habitats. This in part could be due to i) narrower slope gradient in sedimentary grounds than in rocky grounds, and therefore a key role of slope above depth in hard grounds; ii) higher patchy distribution of hard bottoms (and their associated assemblages); iii) higher spatial irregularity in the sampling of hard-bottom assemblages due to its own patchy distribution.

Within the North Atlantic a number of studies have examined faunal zonation by depth and there is strong agreement between studies in respect of the depth of faunal boundaries (Howell, 2010). Studies, however, were rather restricted to soft-bottom zones, while studies on e.g. coral assemblages are comparatively scarce (Wareham and Edinger, 2007 in Newfoundland-Labrador; De Mol et al., 2002 review from Porcupine and N Atlantic) and, likely from the aspects cited above, such

analyses do not give zonation patterns. Our results are coherent with faunal boundaries described in the area, but taking into account seamount topography as an additional key factor, especially the changes in slope gradient in the bank break (at about 1500 m in GB) for hard-bottom assemblages. There is a consistent reporting of faunal boundaries at 500–700 m, 1000–1400 m, 1600–2000 m and 2500–3000 m (Gage, 1986; Gage et al., 2000; Howell et al., 2002; Olabarria, 2005; Howell, 2010; Cartes et al., 2014). The habitat of Ophiacanthidae and *Flabellum* (SS, 750–780 m) would correspond with the upper slope zone in the continental shelf, which ranges from the shelf slope break (depth of the seasonal thermocline) in the continent to the first faunal discontinuity, which in this region lies at approximately 750 m. The 750 m boundary is associated with the top of the permanent thermocline, the

10 °C isotherm, and the boundary between ENAW and MOW. On the GB at 780 m the SS habitat is substituted by the SSrf habitat (*Lophelia* and *Madrepora* reef on sands, 780–1000 m). This zone, defined in the continent as the upper bathyal zone, ranges from the first faunal discontinuity to the second, which in this region is broadly defined as 750–1100 m (Howell et al., 2010). The 1100 m boundary is associated with changes in slope gradient, current speed, and food supply, on the GB corresponding with the beginning of the summit break.

At the limit between the upper and the mid bathyal zone, the BBS habitat, bank break typified by *Cidaris* and *Thenia*, is characterised by a different relief, with a higher presence of rocky outcrops and a higher slope. This habitat is located in the area of influence of the core of MOW (Iorga and Lozier, 1999; Cartes et al., 2014). Christiansen (2010) described as a community of *Thenia muricata* and *Geodia* spp. appears where the seamount slope inclination increases.

FS habitat located on the plain flanks beside GB and typified by the holothurians *Benthogone rosea*, correspond with the mid bathyal zone, which in this region lies between 1100 and 1800 m. The 1800 m boundary is associated with the bottom of the permanent thermocline, the 4 °C isotherm and the boundary between MOW and LSW and possible changes in food supply (Howell et al., 2010). Deeper than our range of study, the lower bathyal zone, between 1800 and 2700 m, and the abyssal zone ranging from 2700 to 5000 m are found. Cartes et al. (2014) defined two assemblages of decapods crustaceans on the GB, mainly on sedimentary areas, at 700–1400 and 1500–1800 m, corresponding with the summit and the flanks and with different water masses.

4.2. Habitat characterisation

The abundance of vulnerable habitats in the GB is directly related to seamount environment. Typically, these areas are associated with strong current regimes (Genin et al., 1986; Herring, 2002). The combination of slopes and strong currents increase the presence of hard substrate. Because of the rarity of hard substrate in the deep sea, areas of high vertical relief often harbour abundant communities of benthic organisms, including deep-water corals (i.a. Herring, 2002; Rogers et al., 2007). On the GB, there is a clear link among the presence of vulnerable habitats and higher slopes (bank break habitats). Seamount conditions favour the presence of sessile vulnerable filter-feeder species such as corals and sponges (Auster et al., 2005; Samadi et al., 2007; Rowden et al., 2010). Sessile filter-feeders rely on currents to provide nutrition as well as to remove sediments that may smother them. Consequently, they are usually associated with strong current velocities or unique current patterns such as recirculation gyres, which in turn are indicative of increased concentrations of particles (Moore and Bullis, 1960; Tendal, 1992). GB vulnerable habitats are typified by sessile filter feeders or zooplankton-carnivore species, such as cold-water corals, bamboo and black corals, gorgonians and large sponges which are located in areas of stronger hydrodynamism such as megaripple sandy areas of the summit and southern slope facing northward currents (Prieto et al., 2013).

Areas of steep relief are known to contribute in enhancing zooplankton aggregations, resulting in zooplankton being the main compartment supporting trophic webs over seamounts (Genin and Dower, 2007). The collision of water masses with seamount walls/slopes is the main argument for the bottom trapping hypothesis, used to explain zooplankton biomass increase and trophic enrichment, e.g. of re-suspended particles over seamounts (Genin and Dower, 2007). On the GB, the highest near-bottom zooplankton biomass (4.3 g/1000 m³), ca. 5 times > than the average on the rest of the bank (Papiol et al., 2014), was found in a planktonic net haul performed in parallel to a vertical wall (at 42°27.36' N- 11°53.84' W, in the southern slope).

Cold-water corals appear on the GB in three different typologies: as reef in sedimentary areas (SSrf; as in Wienberg et al., 2008; De Mol et al., 2002;), as reef in rocky areas (Rucabado Bank: RB; as in Wilson,

1979; Frederiksen et al., 1992) and as discrete colonies in coral gardens together with gorgonians, bamboo and black corals (BBR-S; as in Gubbay, 2003; Howell et al., 2010; Braga-Henriques et al., 2013). Sánchez et al. (2014) described that the presence of living cold-water reefs is directly related to a high-energy environment at depths between 700 and 1200 m, between the lower boundary of ENACW and the core of MOW, in a depth range that matches the water density range $\sigma_\theta = 27.35\text{--}27.65 \text{ kg m}^{-3}$ that has been identified as the limit range for cold-water coral distribution in the North Atlantic (Davies et al., 2005).

Among hard-bottom assemblages, the lowest diversity was at SPR, at the summit, increasing at the slopes, at BBR-N and especially at BBR-S and RB, where the highest diversity was recorded. BBR-S and RB are characterised by the presence of cold-water corals which form complex three-dimensional structures that support a diverse macro- and mega-faunal community, by enhancing habitat complexity and heterogeneity, and providing attachment substrate, shelter, feeding, spawning and nursery areas for other species (Henry and Roberts, 2007; Buhl-Mortensen et al., 2010). More complex assemblages, BBR-N and especially BBR-S/RB, are located at depths within or closer to the influence of Labrador Sea Water (LSW, at 1300 to ca. 1700 m; Yashayaev and Loder, 2009). The enrichment of these deep assemblages of sessile hard-bottom corals and sponges at BBR-N and BBR-S/RB could be made by this northern “Labrador influence”, as is suggested for decapod crustacean assemblages (Cartes et al., 2014). At bank break rock habitats (BBR-N and BBR-S), *Acanella arbuscula*, absent at shallower assemblages, appeared as dominant species only at BBR-N, so at north face of GB, appearing also as a non-dominant species at BBR-S and at the sedimentary assemblages BBS and FS. *Acanella arbuscula* is a dominant coral in Labrador-Newfoundland (Wareham and Edinger, 2007), an area with high levels of coral diversity (28 species) and the highest species richness ($S = 16$) at 200–500 m. *Acanella arbuscula* would not be the only example of this enrichment. Other species (e.g. *Anthothela grandiflora*) were also found at the Labrador shelf-slope (Wareham and Edinger, 2007). These species would cohabit at GB with colonies of cold-water corals (*Lophelia pertusa*, *Madrepora oculata*) and other gorgonians, widely distributed across reefs in different areas of the North Atlantic Ocean (Wilson, 1979; Frederiksen et al., 1992; Sherwood et al., 2005; Sánchez et al., 2008) at ca. 300–800 m, including GB (Duineveld et al., 2004; León et al., 2010; Somoza et al., 2014).

Summit plain rock assemblages, outside these enriched slope areas, are less structurally complex and diverse than slope ones, and are characterised by sparse groups of gorgonians and black corals. Wienberg et al. (2008) described a Rockall Bank assemblage dominated by a fauna similar than SPR in GB: gorgonians (*Acanthogorgia armata*) and antipatharian corals (including *Parantipathes* sp., *Bathypathes* sp., *Stichopathes* sp., and *Leiopathes* sp.).

The other main group of vulnerable sessile filter-feeder, the sponges, is dominant in assemblages found close to the steep escarpment of the bank break, both in sedimentary (BBS) and rocky (BBR-N, BBR-S), in agreement with previous seamount studies (Christiansen, 2010; Davies et al., 2015) and in the same enriched environments described above where cold-water corals are also dominant.

Besides coral and sponges, echinoderms were revealed as one of the key taxa typifying habitats on the GB. Several habitats similar to the SS habitat (summit sands characterised by Ophiacanthidae), at similar depths have been described in literature. Metaxas and Giffin (2004) described dense beds of Ophiacanthidae (*Ophiacantha abyssicola*), in sedimentary areas of Nova Scotia, in depths similar to the Ophiacanthidae habitat on GB. Cherbonnier and Sibuet (1972) and Gage (1983) described the bathymetrical preference of the species *Ophiomyces grandis* for 200–700 m range. The spatial distribution of GB ophiuroids habitat in an area of strong currents (presence of megaripples) could be a consequence of trophic-hydrographic drivers. Dense populations of *Ophiocten gracilis* able to intercept particles from near-bed flow, coincides with the slope current and largely disappeared at 1000 m (Lamont and Gage, 1998; Gage et al., 2000).

The link between habitat distribution and oceanographic circulation is obvious, since most seamount habitats are typified by filter feeders (Samadi et al., 2007; Rogers et al., 2007; Rowden et al., 2010). Data from three surveys has revealed oceanographic patterns over the GB seamount. Waiting for a specific study on topographical effects, a link with habitat complexity and currents seems probable, since habitats oriented south and southwest show higher complexity probably due to predominance of northward currents. Prieto et al. (2013) stated Mediterranean water (MOW), spreading from the Strait of Gibraltar, flows northwards along the continental slope sometimes showing a detachment contouring the southwest Galicia Bank.

4.3. GB habitats classification

Several problems appeared when deep-sea habitats described in this paper were translated to EUNIS level 4–6 since there are no direct equivalencies. Only cold-water coral reef among GB habitat has a coherent counterpart in EUNIS. Regarding HD, only RB, BBR-S and BBR-N (coral reef, coral gardens and sponges on slope rock) and SSrf (cold-water coral reef on sands) fulfill the conditions required to be included in the habitat type 1170 “Reefs”. The low density of gorgonians and black corals in the summit plain rock is the reason to not include this habitat in the HD 1170. GB habitats can be also included in the OSPAR list of threatened habitats (OSPAR, 2008). Cold-water coral reefs located in the Rucabado Bank and in the summit sands are clearly habitats belonging to the *Lophelia* reefs and/or Carbonate mounds types. Ophiacanthidae and *Flabellum* habitat and the rocky habitats with gorgonians, black and bamboo corals can be included in the type Coral Gardens. The habitat typified by *Thenaea* can be included in the OSPAR list as a Deep-sea Sponge aggregation, since this sponge is included in the list of species frequently reported from Sponge Grounds in the NEA (Christiansen, 2010), as a component species of the aggregations characterised by geodids or constituting its own characteristic habitat.

The hierarchical cold-water corals biotope classification scheme proposed by Davies et al. (2017) is adequate to classify the GB biotopes formed by corals. The only exception is the absence in this scheme of hard substrate habitats formed by the bamboo coral *Acanella arbuscula*, common in the GB rocky grounds.

4.4. GB environmental values as a SAC

The GB proposal as a SAC was based on the presence of vulnerable habitats and species. Coral assemblages found on seamounts and island slopes showed high taxonomic richness at both species and family levels, indicating that the benthic habitats of the seamounts harbour numerous species currently recognised as key indicators of vulnerable marine ecosystems (OSPAR, 2008; FAO, 2009). We have already described how seamount conditions in general (Auster et al., 2005; Rogers et al., 2007; Rowden et al., 2010) and GB in particular (Somoza et al., 2014) favour the existence of habitats built by sessile vulnerable filter-feeder species such as corals and sponges.

Some habitats described in this paper are formed by slow-growing and fragile species and hence are potentially vulnerable, but are not included in HD Annex since they are not “reefs” (e.g. Ophiacanthidae and *Flabellum*, *Cidaris* and *Thenaea*). The conservation of vulnerable habitats outside rocky areas should be taken into account in the future.

Another value is the singularity of GB habitats and communities. In the northeastern Atlantic, Galicia Bank is considered a coastal seamount, together with the Ampere, Gorringe, Josephine and Seine banks, in contrast with oceanic seamounts, including the Atlantis, Hyeres, Irving, Meteor and Plato banks located offshore (Gofas, 2007; Surugiu et al., 2008). Nevertheless, some results on inter-seamount faunal similarity highlight the separation of GB due to its isolated northern position and deep plateau (Surugiu et al., 2008). A key factor that controls the occurrence and particularly the abundance of benthic animals is the water depth at the seamount's summit (Clark et al., 2011;

Tempera et al., 2012). Seamounts that rise to approximately 1500 m in depth have much higher densities of faunal coverage than deeper seamounts (Harris, 2012). In the case of GB, a combination of factors: depth, substrate type and the influence of water masses determine hard and soft bottom communities, which in the case of corals, it seems enrichment from northern water masses (LSW) arriving to GB and a possible zooplankton biomass increase at vertical-steep walls by “bottom trapping” (Genin and Dower, 2007) can explain higher diversity at slope rocky breaks.

The combination of clustering and modeling used in this paper represents a useful and feasible technique to generate habitat maps at the EUNIS level 4–6 in a standardised way which follows robust scientific criteria to correctly answer HD and MSFD directives. The results of this paper were the basis for the proposal of GB as a SAC and will be useful to state the environmental values which have to be taken into account in the future GB management plan to protect its ecosystem function and biodiversity, and its significance as a relatively unexploited example of a seamount within the Natura 2000 network of marine protected areas in the NE Atlantic.

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Appendix A. Supplementary material

Supplementary data associated with this article can be found in the online version at <http://dx.doi.org/10.1016/j.dsr.2017.06.003>.

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